SEQUENCE LISTING

<110> FIEBIG, HELMUT NANDY, ANDREAS CROMWELL, OLIVER
<120> DNA SEQUENCE, AND RECOMBINANT PREPARATION OF GROUP 4 MAJOR ALLERGENS FROM CEREALS
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<140> 10/583,089 <141> 2006-06-15
<150> PCT/EP04/013664 <151> 2004-12-01
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ctc gga tgc ctc atg aag gag ata ccg gcc cgc ctc ctc tac gcc aag 144 Leu Gly Cys Leu Met Lys Glu Ile Pro Ala Arg Leu Leu Tyr Ala Lys 35 40 45
agc tcg cct gac tac ccc acc gtg ctg gcg cag acc atc agg aac tcg 192 Ser Ser Pro Asp Tyr Pro Thr Val Leu Ala Gln Thr Ile Arg Asn Ser 50 55 60
cgg tgg tcg tcg ccg cag aac gtg aag ccg atc tac atc atc acc ccc 240 Arg Trp Ser Ser Pro Gln Asn Val Lys Pro Ile Tyr Ile Ile Thr Pro 65 70 75 80
acc aac gcc tcg cac atc cag tcc gcg gtg gtg tgc ggc cgc cgg cac Thr Asn Ala Ser His Ile Gln Ser Ala Val Val Cys Gly Arg Arg His 85 90 95

				_	gtg Val		_				_				_	336
_					aaa Lys			_		_	_	_	_			384
					tcg Ser	_	_			_	_	_			_	432
_				_	ctc Leu 150								_	_		480
					ttc Phe											528
				_	ggc Gly					_	_	_	_			576
		_	_		aac Asn	_		_	_	_		-	_			624
					aag Lys											672
					gga Gly 230		_				_	_	_		_	720
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_	_			_	ggc Gly	_		_		_		_		_	_	816
					ccc Pro											864
					ttc Phe											912
					agc Ser 310											960

· ·

_		tgc Cys			_				_							1008
		aag Lys	_			_	_									1056
		ttc Phe 355														1104
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		gtc Val					_			_				_	_	1296
		agc Ser 435														1344
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	_	aag Lys			_							_			_	1488
		gtg Val														1536
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Ser Ser Pro Asp Tyr Pro Thr Val Leu Ala Gln Thr Ile Arg Asn Ser 50 55 60

Arg Trp Ser Ser Pro Gln Asn Val Lys Pro Ile Tyr Ile Ile Thr Pro 65 70 75 80

Thr Asn Ala Ser His Ile Gln Ser Ala Val Val Cys Gly Arg Arg His
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Gly Ile Arg Leu Arg Val Arg Ser Gly Gly His Asp Tyr Glu Gly Leu 100 105 110

Ser Tyr Arg Ser Glu Lys Pro Glu Thr Phe Ala Val Val Asp Leu Asn 115 120 125

Lys Met Arg Ala Val Ser Val Asp Gly Tyr Ala Arg Thr Ala Trp Val 130 135 140

Ser Pro Val Leu Ala Phe Pro Ala Gly Val Cys Pro Ser Ile Gly Val
165 170 175

Gly Gly Asn Phe Ala Gly Gly Gly Phe Gly Met Leu Leu Arg Lys Tyr 180 185 190

Gly Ile Ala Ala Glu Asn Val Ile Asp Val Lys Val Val Asp Pro Asn 195 200 205

Gly Lys Leu Leu Asp Lys Ser Ser Met Ser Ala Asp His Phe Trp Ala 210 215 220

Val Arg Gly Gly Gly Glu Ser Phe Gly Ile Val Val Ser Trp Gln 225 230 235 240

Val Lys Leu Leu Pro Val Pro Pro Thr Val Thr Val Leu Lys Ile Pro 245 250 255

Lys Thr Val Gln Glu Gly Ala Ile Asp Leu Val Asn Lys Trp Gln Leu 260 265 270

Val Gly Pro Ala Leu Pro Gly Asp Leu Met Ile Arg Ile Ile Leu Ala 275 280 285

Gly Asn Ser Ala Thr Phe Glu Ala Met Tyr Leu Gly Thr Cys Ser Thr 290 295 300

Leu Thr Pro Leu Met Ser Ser Lys Phe Pro Glu Leu Gly Met Asn Pro 305 310 315 320

Ser His Cys Asn Glu Met Ser Trp Ile Lys Ser Ile Pro Phe Ile His 325 330 335

Leu Gly Lys Gln Asn Leu Asp Asp Leu Leu Asn Arg Asn Asn Thr Phe 340 345 350

Lys Pro Phe Ala Glu Tyr Lys Ser Asp Tyr Val Tyr Gln Pro Phe Pro 355 360 365

Lys Pro Val Trp Glu Gln Ile Phe Gly Trp Leu Val Lys Pro Gly Ala 370 375 380

Gly Ile Met Ile Met Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro 385 390 395 400

Glu Ala Ala Thr Pro Phe Pro His Arg Gln Gly Val Leu Phe Asn Ile 405 410 415

Gln Tyr Val Asn Tyr Trp Phe Ala Glu Ser Ala Gly Ala Ala Pro Leu 420 425 430

Gln Trp Ser Lys Asp Ile Tyr Lys Phe Met Glu Pro Tyr Val Ser Lys 435 440 445

Asn Pro Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp Leu Gly Arg 450 450 460

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		acc Thr														288
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		aag Lys	_		_			_	_		_	_	_	_		432
	-	gac Asp				_							_		_	480
aag Lys	aac Asn	agc Ser	ccc Pro	gtg Val 165	ctc Leu	gcg Ala	ttc Phe	ccg Pro	gcc Ala 170	ggc Gly	gtt Val	tgc Cys	ccg Pro	acc Thr 175	att Ile	528
		ggc Gly														576
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		ggc Gly														672

					ggc Gly 230											720
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		_	_		caa Gln	-		_	_					_		816
					gcc Ala											864
_					gcc Ala	_			_	_		_			_	912
	_	_		_	ctg Leu 310	_	_	_	_						_	960
	_	_		_	aac Asn		_				_		_			1008
Ile	His	Leu	Gly 340	Lys	cag Gln	Ala	Thr	Leu 345	Ser	Āsp	Leu	Leu	Asn 350	Arg	Asn	1056
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Pro	Val 370	Pro	Lys	Pro	gtc Val	Trp 375	Ala	Gln	Ile	Phe	Val 380	Trp	Leu	Val	Lys	1152
Pro 385	Gly	Ala	Gly	Ile	atg Met 390	Val	Met	Asp	Pro	Tyr 395	Gly	Ala	Ala	Ile	Ser 400	1200
Ala	Thr	Pro	Glu	Ala 405	gcc Ala	Thr	Pro	Phe	Pro 410	His	Arg	Lys	Asp	Val 415	Leu	1248
Phe	Asn	Ile	Gln 420	Tyr	gtc Val	Asn	Tyr	Trp 425	Phe	Asp	Glu	Ala	Gly 430	Gly	Ala	1296
					agc Ser											1344

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ctc ggc agg aac gag gtg gtc aac gac atc tcc acc tat gcc agc ggc Leu Gly Arg Asn Glu Val Val Asn Asp Ile Ser Thr Tyr Ala Ser Gly 465 470 475 480	1440
aag gtc tgg ggc gag aag tac ttc aag ggc aac ttc caa agg ctc gcc Lys Val Trp Gly Glu Lys Tyr Phe Lys Gly Asn Phe Gln Arg Leu Ala 485 490 495	1488
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Asn Ser Arg Ala Phe Ala Leu Val Pro Leu Leu Ile Cys Val Leu Ser 1 5 10 15 Cys His Ala Ala Val Ser Tyr Ala Ala Ala Pro Val Pro Ala Lys Glu 20 25 30 Asp Phe Phe Gly Cys Leu Val Lys Glu Ile Pro Ala Arg Leu Leu Tyr	
Asn Ser Arg Ala Phe Ala Leu Val Pro Leu Leu Ile Cys Val Leu Ser 15 Cys His Ala Ala Val Ser Tyr Ala Ala Ala Pro Val Pro Ala Lys Glu 25 Asp Phe Phe Gly Cys Leu Val Lys Glu Ile Pro Ala Arg Leu Leu Tyr 35 Ala Lys Ser Ser Pro Ala Phe Pro Thr Val Leu Ala Gln Thr Ile Arg	
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Asn Ser Arg Ala Phe Ala Leu Val Pro Leu Leu Ile Cys Val Leu Ser 15 Cys His Ala Ala Val Ser Tyr Ala Ala Ala Pro Val Pro Ala Lys Glu 25 Asp Phe Phe Gly Cys Leu Val Lys Glu Ile Pro Ala Arg Leu Leu Tyr 35 Ala Lys Ser Ser Pro Ala Phe Pro Thr Val Leu Ala Gln Thr Ile Arg 50 Asn Ser Arg Trp Ser Ser Pro Gln Ser Val Lys Pro Leu Tyr Ile Ile 65 Thr Pro Thr Asn Ala Ser His Ile Gln Ser Ala Val Val Cys Gly Arg	
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Trp Val Asp Ser Gly Ala Gln Leu Gly Glu Leu Tyr Tyr Ala Ile Ala 145 150 155 160

Lys Asn Ser Pro Val Leu Ala Phe Pro Ala Gly Val Cys Pro Thr Ile 165 170 175

Gly Val Gly Gly Asn Phe Ala Gly Gly Gly Phe Gly Met Leu Leu Arg 180 185 190

Lys Tyr Gly Ile Ala Ala Glu Asn Val Ile Asp Val Lys Val Val Asp 195 200 205

Ala Asn Gly Thr Leu Leu Asp Lys Ser Ser Met Ser Ala Asp His Phe 210 215 220

Trp Ala Val Arg Gly Gly Gly Glu Ser Phe Gly Ile Val Val Ser 225 230 235 240

Trp Gln Val Lys Leu Leu Pro Val Pro Pro Thr Val Thr Val Phe Lys
245 250 255

Ile Pro Lys Thr Val Gln Glu Gly Ala Val Glu Leu Ile Asn Lys Trp
260 265 270

Gln Leu Val Ala Pro Ala Leu Pro Asp Asp Leu Met Ile Arg Ile Ile 275 280 285

Ala Phe Gly Gly Thr Ala Lys Phe Glu Ala Met Tyr Leu Gly Thr Cys 290 295 300

Lys Ala Leu Thr Pro Leu Met Ser Ser Arg Phe Pro Glu Leu Gly Met 305 310 315 320

Asn Ala Ser His Cys Asn Glu Met Pro Trp Ile Lys Ser Val Pro Phe 325 330 335

Ile His Leu Gly Lys Gln Ala Thr Leu Ser Asp Leu Leu Asn Arg Asn 340 345 350

Asn Thr Phe Lys Pro Phe Ala Glu Tyr Lys Ser Asp Tyr Val Tyr Gln 355 360 365

Pro Val Pro Lys Pro Val Trp Ala Gln Ile Phe Val Trp Leu Val Lys 370 375 380

Pro Gly Ala Gly Ile Met Val Met Asp Pro Tyr Gly Ala Ala Ile Ser 385 390 395 400

Ala Thr Pro Glu Ala Ala Thr Pro Phe Pro His Arg Lys Asp Val Leu 405 410 415

Phe Asn Ile Gln Tyr Val Asn Tyr Trp Phe Asp Glu Ala Gly Gly Ala 420 425 430

Ala Pro Leu Gln Trp Ser Lys Asp Met Tyr Arg Phe Met Glu Pro Tyr 435 440 445

Val Ser Lys Asn Pro Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp

Leu Gly Arg Asn Glu Val Val Asn Asp Ile Ser Thr Tyr Ala Ser Gly 475 Lys Val Trp Gly Glu Lys Tyr Phe Lys Gly Asn Phe Gln Arg Leu Ala Ile Thr Lys Gly Lys Val Asp Pro Gln Asp Tyr Phe Arg Asn Glu Gln 505 Ser Ile Pro Pro Leu Leu Gly Lys <210> 5 <211> 1608 <212> DNA <213> Hordeum vulgare <220> <221> CDS <222> (1)..(1557) <400> 5 age teg agg gee tte get etg gtg etc etc etc tge gee ttg tee tge 48 Ser Ser Arg Ala Phe Ala Leu Val Leu Leu Cys Ala Leu Ser Cys cac cac get gec gtc tee tee geg cag gtg eeg gee aag gae gae tte His His Ala Ala Val Ser Ser Ala Gln Val Pro Ala Lys Asp Asp Phe 20 25 ctg gga tgc ctc gtg aag gag ata ccg gcc cgc ctc ctc ttc gcc aag Leu Gly Cys Leu Val Lys Glu Ile Pro Ala Arg Leu Leu Phe Ala Lys age teg cet gee tte eee gee gte etg gag cag ace ate agg aac teg 192 Ser Ser Pro Ala Phe Pro Ala Val Leu Glu Gln Thr Ile Arg Asn Ser cgg tgg tcg tcg ccg cag aac gtg aag ccg ctc tac atc acc ccc 240 Arg Trp Ser Ser Pro Gln Asn Val Lys Pro Leu Tyr Ile Ile Thr Pro ace ace tee cac ate cag tet get gtg gtg tge gge ege egg cae 288 Thr Asn Thr Ser His Ile Gln Ser Ala Val Val Cys Gly Arg Arg His

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105

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tcg tac Ser Ty				_					_	_	_	_			384
aag ato Lys Met 130	Arg														432
gac tco Asp Ser 145			_							-		_	_		480
agc ccc Ser Pro															528
ggt ggd Gly Gly															576
ggc ato Gly Ile															624
ggc aag Gly Lys 210	Leu														672
gtc agg Val Arg 225						_				_	_	_		_	720
gtg aag Val Lys			-									_			768
aag aca Lys Thi															816
gtc gcg Val Ala															864
ggg gad Gly Asp 290	Lys	gcg Ala	acg Thr	ttc Phe	gag Glu 295	gcc Ala	atg Met	tac Tyr	ctg Leu	ggc Gly 300	acc Thr	tgc Cys	aaa Lys	acc Thr	912
ctg acc Leu Thr 305															960
tcg cac Ser His															1008

		_	_	_		_	_	-						aac Asn		1056
				_	_		_	_	_		-		_	ccc Pro	_	1104
														ccc Pro		1152
			_	_	_	_				_			_	gcc Ala		1200
														ttc Phe 415		1248
	_		-					_		_	_		_	gcg Ala	_	1296
														gtg Val		1344
							_				-		_	ctc Leu		1392 ·
						_					_	-		aag Lys		1440
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- Arg Trp Ser Ser Pro Gln Asn Val Lys Pro Leu Tyr Ile Ile Thr Pro 65 70 75 80
- Thr Asn Thr Ser His Ile Gln Ser Ala Val Val Cys Gly Arg Arg His
 85 90 95
- Gly Val Arg Leu Arg Val Arg Ser Gly Gly His Asp Tyr Glu Gly Leu 100 105 110
- Ser Tyr Arg Ser Glu Arg Pro Glu Ala Phe Ala Val Val Asp Leu Asn 115 120 125
- Lys Met Arg Thr Val Leu Val Asn Glu Lys Ala Arg Thr Ala Trp Val 130 135 140
- Ser Pro Val Leu Ala Phe Pro Ala Gly Val Cys Pro Ser Ile Gly Val
 165 170 175
- Gly Gly Asn Phe Ala Gly Gly Gly Phe Gly Met Leu Leu Arg Lys Tyr 180 185 190
- Gly Ile Ala Ala Glu Asn Val Ile Asp Val Lys Leu Val Asp Ala Asn 195 200 205
- Gly Lys Leu Leu Asp Lys Ser Ser Met Ser Pro Asp His Phe Trp Ala 210 215 220
- Val Arg Gly Gly Gly Glu Ser Phe Gly Ile Val Val Ser Trp Gln 225 230 235 240
- Val Lys Leu Pro Val Pro Pro Thr Val Thr Val Phe Gln Ile Pro
 245 250 255
- Lys Thr Val Gln Glu Gly Ala Val Asp Leu Ile Asn Lys Trp Gln Leu 260 265 270
- Val Ala Pro Ala Leu Pro Gly Asp Ile Met Ile Arg Ile Ile Ala Met 275 280 285
- Gly Asp Lys Ala Thr Phe Glu Ala Met Tyr Leu Gly Thr Cys Lys Thr 290 295 300

Leu Thr Pro Leu Met Ser Ser Lys Phe Pro Glu Leu Gly Met Asn Pro 305 310 315 320

Ser His Cys Asn Glu Met Pro Trp Ile Lys Ser Ile Pro Phe Ile His 325 330 335

Leu Gly Lys Gln Ala Thr Leu Ala Asp Leu Leu Asn Arg Asn Asn Thr 340 345 350

Phe Lys Pro Phe Ala Glu Tyr Lys Ser Asp Tyr Val Tyr Gln Pro Val 355 360 365

Pro Lys Pro Val Trp Glu Gln Leu Phe Gly Trp Leu Thr Lys Pro Gly 370 380

Ala Gly Ile Met Val Met Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr 385 390 395 400

Pro Glu Ala Ala Thr Pro Phe Pro His Arg Lys Gly Val Leu Phe Asn 405 410 415

Ile Gln Tyr Val Asn Tyr Trp Phe Ala Glu Ala Ala Gly Ala Ala Pro 420 425 430

Leu Gln Trp Ser Lys Asp Ile Tyr Lys Phe Met Glu Pro Phe Val Ser 435 440 445

Lys Asn Pro Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp Leu Gly 450 455 460

Arg Asn Glu Val Val Asn Asp Ile Ser Thr Tyr Ser Ser Gly Lys Val 465 470 475 480

Trp Gly Glu Lys Tyr Phe Lys Gly Asn Phe Gln Arg Leu Ala Ile Thr 485 490 495

Lys Gly Lys Val Asp Pro Gln Asp Tyr Phe Arg Asn Glu Gln Ser Ile 500 505 510

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_	cct Pro 50	-				_	_		_					_		192
	ttg Leu															240
	gcc Ala															288
_	cgc Arg		_	_		-				_				_	_	336
	cgg Arg						_		-	_	_	_			_	384
	cgg Arg 130															432
	ggc Gly															480
	gtg Val															528
	aac Asn															576
	gcc Ala															624
	ctt Leu 210		_	_	_		_	_	_	_				_	_	672
	ggc Gly	_											-			720

				gtg Val 245												768
				ggc Gly												816
				ccc Pro												864
				ttc Phe												912
				agc Ser												960
				atg Met 325												1008
				agc Ser												1056
				gaa Glu												1104
aag Lys	ccc Pro 370	gtg Val	tgg Trp	gag Glu	cag Gln	atc Ile 375	ttc Phe	ggc Gly	tgg Trp	ctc Leu	acg Thr 380	aag Lys	ccc Pro	ggt Gly	gjå aaa	1152
				atg Met												1200
gaa Glu				ccg Pro 405												1248
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aac Asn	ccc Pro 450	agg Arg	cag Gln	gcg Ala	tac Tyr	gcc Ala 455	aac Asn	tac Tyr	agg Arg	gac Asp	att Ile 460	gac Asp	ctc Leu	ggc Gly	agg Arg	1392

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Ser Pro Asp Phe Pro Thr Val Leu Ala Gln Thr Ile Arg Asn Ser Arg 50 55 60	
Trp Leu Ser Pro Gln Asn Val Lys Pro Leu Tyr Ile Ile Thr Pro Thr 65 70 75 80	
Asn Ala Ser His Ile Gln Ser Ala Val Val Cys Gly Arg Arg His Ser 85 90 95	
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Tyr Arg Ser Glu Lys Pro Glu Thr Phe Ala Val Val Asp Leu Asn Lys 115 120 125	
Met Arg Ala Val Leu Ile Asp Gly Tyr Ala Arg Thr Ala Trp Val Glu 130 135 140	
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Pro Val Leu Ala Phe Pro Ala Gly Val Cys Pro Thr Ile Gly Val Gly 165 170 175	

- Gly Asn Phe Ala Gly Gly Phe Gly Met Leu Leu Arg Lys Tyr Gly 180 185 190
- Ile Ala Ala Glu Asn Val Ile Asp Val Lys Val Val Asp Pro Asn Gly 195 200 205
- Lys Leu Leu Asp Lys Ser Ser Met Ser Pro Asp His Phe Trp Ala Val 210 215 220
- Arg Gly Gly Gly Glu Ser Phe Gly Ile Val Val Ser Trp Gln Val 225 230 235 240
- Lys Leu Leu Pro Val Pro Pro Thr Val Thr Val Phe Lys Ile Pro Lys 245 250 255
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 260 265 270
- Gly Pro Ala Leu Pro Gly Asp Leu Met Ile Arg Val Ile Ala Ala Gly
 275 280 285
- Asn Thr Ala Thr Phe Glu Gly Met Tyr Leu Gly Thr Cys Gln Thr Leu 290 295 300
- Thr Pro Leu Met Ser Ser Gln Phe Pro Glu Leu Gly Met Asn Pro Tyr 305 310 315 320
- His Cys Asn Glu Met Pro Trp Ile Lys Ser Ile Pro Phe Ile His Leu 325 330 335
- Gly Lys Glu Ala Ser Leu Val Asp Leu Leu Asn Arg Asn Asn Thr Phe 340 345 350
- Lys Pro Val Trp Glu Gln Ile Phe Gly Trp Leu Thr Lys Pro Gly Gly 370 375 380
- Gly Met Met Ile Met Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro 385 390 395 400
- Glu Ala Ala Thr Pro Phe Pro His Arg Gln Gly Val Leu Phe Asn Ile 405 410 415
- Gln Tyr Val Asn Tyr Trp Phe Ala Glu Ala Ala Ala Ala Pro Leu 420 425 430
- Gln Trp Ser Lys Asp Met Tyr Asn Phe Met Glu Pro Tyr Val Ser Lys 435 440 445
- Asn Pro Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp Leu Gly Arg 450 455 460
- Asn Glu Val Val Asn Asp Ile Ser Thr Tyr Ser Ser Gly Lys Val Trp 465 470 475 480

Gly Glu Lys Tyr Phe Lys Gly Asn Phe Gln Arg Leu Ala Ile Thr Lys 485 490 495

Gly Lys Val Asp Pro Gln Asp Tyr Phe Arg Asn Glu Gln Ser Ile Pro 500 505 510

Pro Leu Leu Glu Lys Tyr 515

130

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	gtg Val				_	_		_	_	_				_		528
	aac Asn								_	_	_	_	_			576
	gcc Ala	_			_		_	_	_		_	_		_		624
	ctg Leu 210															672
	ggc					_				_	_	_		_		720
_	ctc Leu	_						_		_		_			_	768
	gtg Val				_	_	_		_		_		_	_	_	816
	ccg Pro															864
	acg Thr 290															912
	ccg Pro															960
cac His	tgc Cys	aac Asn	gag Glu	atg Met 325	ccc Pro	tgg Trp	atc Ile	aag Lys	tcc Ser 330	gtc Val	ccc Pro	ttc Phe	atc Ile	cac His 335	ctc Leu	1008
ggc Gly	aaa Lys	cag Gln	gct Ala 340	ggc	ctg Leu	gac Asp	gac Asp	ctc Leu 345	ctc Leu	aac Asn	cgg Arg	aac Asn	aac Asn 350	acc Thr	ttc Phe	1056
	ccc Pro															1104

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ggg atc atg atc atg gac ccc tac ggc gcc acc atc agc gcc acc ccc Gly Ile Met Ile Met Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro 385 390 395 400	1200									
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cag tgg agc aag gac att tac aat ttc atg gag ccg tac gtg agc aag Gln Trp Ser Lys Asp Ile Tyr Asn Phe Met Glu Pro Tyr Val Ser Lys 435 440 445	1344									
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- Trp Ser Thr Gln Gln Asn Val Lys Pro Leu Tyr Ile Ile Thr Pro Thr 65 70 75 80
- Asn Ala Ser His Ile Gln Ser Ala Val Val Cys Gly Arg Arg His Gly 85 90 95
- Val Arg Leu Arg Val Arg Ser Gly Gly His Asp Tyr Glu Gly Leu Ser 100 105 110
- Tyr Arg Ser Glu Lys Pro Glu Thr Phe Ala Val Val Asp Leu Asn Lys 115 120 125
- Met Arg Ala Val Val Val Asp Gly Tyr Ala Arg Thr Ala Trp Val Glu 130 135 140
- Ser Gly Ala Gln Leu Gly Glu Leu Tyr Tyr Ala Ile Ala Lys Asn Ser 145 150 155 160
- Pro Val Leu Ala Phe Pro Ala Gly Val Cys Pro Ser Ile Gly Val Gly
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- Gly Asn Phe Ala Gly Gly Gly Phe Gly Met Leu Leu Arg Lys Tyr Gly 180 185 190
- Ile Ala Ala Glu Asn Val Ile Asp Val Lys Val Val Asp Pro Asp Gly
 195 200 205
- Lys Leu Leu Asp Lys Ser Ser Met Ser Ala Asp His Phe Trp Ala Val 210 215 220
- Arg Gly Gly Gly Glu Ser Phe Gly Ile Val Val Ser Trp Gln Vai 225 230 235 240
- Lys Leu Met Pro Val Pro Pro Thr Val Thr Val Phe Lys Ile Pro Lys 245 250 255
- Thr Val Gln Glu Gly Ala Val Asp Leu Val Asn Lys Trp Gln Leu Val 260 265 270
- Gly Pro Ala Leu Pro Gly Asp Leu Met Ile Arg Val Ile Ala Ala Gly 275 280 285
- Asn Thr Ala Thr Phe Glu Ala Leu Tyr Leu Gly Thr Cys Lys Thr Leu 290 295 300
- Thr Pro Leu Met Ser Ser Gln Phe Pro Glu Leu Gly Met Asn Pro Tyr 305 310 315 320
- His Cys Asn Glu Met Pro Trp Ile Lys Ser Val Pro Phe Ile His Leu 325 330 335
- Gly Lys Gln Ala Gly Leu Asp Asp Leu Leu Asn Arg Asn Asn Thr Phe 340 345 350

Lys Pro Phe Ala Glu Tyr Lys Ser Asp Tyr Val Tyr Gln Pro Phe Pro Lys Pro Val Trp Glu Gln Ile Phe Gly Trp Leu Ala Lys Pro Gly Ala Gly Ile Met Ile Met Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro Glu Ala Ala Thr Pro Phe Pro His Arg Gln Gly Val Leu Phe Asn Ile 410 Gln Tyr Val Asn Tyr Trp Phe Ala Glu Pro Ala Gly Ala Ala Pro Leu Gln Trp Ser Lys Asp Ile Tyr Asn Phe Met Glu Pro Tyr Val Ser Lys Asn Pro Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp Leu Gly Arg Asn Glu Val Val Asn Asp Ile Ser Thr Tyr Ser Ser Gly Lys Val Trp 465 470 Gly Glu Lys Tyr Phe Lys Ser Asn Phe Gln Arg Leu Ala Ile Thr Lys Gly Lys Val Asp Pro Gln Asp Tyr Phe Arg Asn Glu Gln Ser Ile Pro 500 505 Pro Leu Ile Glu Lys Tyr 515 <210> 11 <211> 1503 <212> DNA <213> Phleum pratense <220> <221> CDS <222> (1)..(1503) <400> 11 tac ttc ccg ccg ccg gct gct aaa gaa gac ttc ctg ggt tgc ctg gtt 48 Tyr Phe Pro Pro Pro Ala Ala Lys Glu Asp Phe Leu Gly Cys Leu Val

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40

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				gtg Val 70									240
				cac His	_				_			_	288
		_		gcc Ala	-	_	_		_	_		 ~ ~	336
				gcc Ala									384
				tac Tyr									432
				tgc Cys 150									480
				atg Met									528
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				gac Asp									624
				atc Ile									672
				aca Thr 230									720
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					tac Tyr											1152
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					aac Asn											1296
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1503

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Pro Ser Val Leu Gly Gln Thr Ile Arg Asn Ser Arg Trp Ser Ser Pro 35 40 45

Asp Asn Val Lys Pro Leu Tyr Ile Ile Thr Pro Thr Asn Val Ser His 50 55 60

Ile Gln Ser Ala Val Val Cys Gly Arg Arg His Ser Val Arg Ile Arg
65 70 75 80

Val Arg Ser Gly Gly His Asp Tyr Glu Gly Leu Ser Tyr Arg Ser Leu 85 90 95

Gln Pro Glu Thr Phe Ala Val Val Asp Leu Asn Lys Met Arg Ala Val 100 105 110

Trp Val Asp Gly Lys Ala Arg Thr Ala Trp Val Asp Ser Gly Ala Gln
115 120 125

Leu Gly Glu Leu Tyr Tyr Ala Ile Tyr Lys Ala Ser Pro Thr Leu Ala 130 135 140

Phe Pro Ala Gly Val Cys Pro Thr Ile Gly Val Gly Asn Phe Ala 145 150 155 160

Gly Gly Phe Gly Met Leu Leu Arg Lys Tyr Gly Ile Ala Ala Glu 165 170 175

Asn Val Ile Asp Val Lys Leu Val Asp Ala Asn Gly Lys Leu His Asp 180 185 190

Lys Lys Ser Met Gly Asp Asp His Phe Trp Ala Val Arg Gly Gly 195 200 205

Gly Glu Ser Phe Gly Ile Val Val Ala Trp Gln Val Lys Leu Leu Pro 210 215 220

Val Pro Pro Thr Val Thr Ile Phe Lys Ile Ser Lys Thr Val Ser Glu 225 230 235 240

Gly Ala Val Asp Ile Ile Asn Lys Trp Gln Val Val Ala Pro Gln Leu 245 250 255 Pro Ala Asp Leu Met Ile Arg Ile Ile Ala Gln Gly Pro Lys Ala Thr 260 265 270

Phe Glu Ala Met Tyr Leu Gly Thr Cys Lys Thr Leu Thr Pro Leu Met 275 280 285

Ser Ser Lys Phe Pro Glu Leu Gly Met Asn Pro Ser His Cys Asn Glu 290 295 300

Met Ser Trp Ile Gln Ser Ile Pro Phe Val His Leu Gly His Arg Asp 305 310 315 320

Ala Leu Glu Asp Asp Leu Leu Asn Arg Asn Asn Ser Phe Lys Pro Phe 325 330 335

Ala Glu Tyr Lys Ser Asp Tyr Val Tyr Gln Pro Phe Pro Lys Thr Val 340 345 350

Trp Glu Gln Ile Leu Asn Thr Trp Leu Val Lys Pro Gly Ala Gly Ile 355 360 365

Met Ile Phe Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro Glu Ser 370 375 380

Ala Thr Pro Phe Pro His Arg Lys Gly Val Leu Phe Asn Ile Gln Tyr 385 390 395 400

Val Asn Tyr Trp Phe Ala Pro Gly Ala Ala Ala Pro Leu Ser Trp
405 410 415

Ser Lys Asp Ile Tyr Asn Tyr Met Glu Pro Tyr Val Ser Lys Asn Pro 420 425 430

Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp Leu Gly Arg Asn Glu
435 440 445

Val Val Asn Asp Val Ser Thr Tyr Ala Ser Gly Lys Val Trp Gly Gln 450 455 460

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